

saug66I
nlaIV
avaII
ppumI
ai

1 AATTCAAGCCAGAGAGCCCTGCCATTTCGTGTGGGCTCAGAGTCCCTACTGCTCAGCCCCCTTCCCTCCGCAAGGCCAATGACCCGGGAGTCCCTTTT
TTAAGTTCGGGCTCTCGGACGGTAAAGACACCCGAGTCCAGGGATGACGATCGGGAGAGGAGGCCGTTCCGCTGTACTTGGCCCCCAGGAAAA
-25
101 AGGCACCTGCTCTGCTGGTGGCTGCTCACTGAGGCTCCAGCAGCCCTCCAGCAGCCATCAGGAAACAAGTGGTCTGGCCAAAAAGGGAGTACGTGAACTGACCT
TCCGTGAACCAAGACCAACGACGCTTGACCCGCGAGGAGGCTCGCTGGTGAAGTCCCTTGTTCACCAAGACCCGTTTTCCTTATGTACCTTGACTGGA
-18 ArgHisLeuLeuValLeuGlnLeuAlaLeuLeuProAlaIleThrGlnGlyAsnGlyValIleuGlyLysGlyAspThrValGluLeuThrCys
201 GTACAGCTTCCCAAGAGAAGCATACATTCACCTGGAAAACTCCACACAGATAAAGATTCTGGAAATCAGGGCTCTTCTTAATAAGTCCATC
rsai mboii hinf
17 ThrAlaSerGlnLysLysSerIleGlnPheHisTyrPlysAsnSerAsnGlnIleLysIleLeuGlyAsnGlnGlySerPheLeuThrLysGlyProSer
301 CAAGCTGAATGATCGCGCTGACTCAAGAGAAGACCTTTGGGACCAAGAACTTTCCTGATCATCAGAATCTTAAGATAGAAGACTCAGATTAATTAC
alul dpmi hinf mboii nlaiv bclI hinf mboii
50 LysLeuAsnAspArgAlaAspSerArgArgSerLeuTyrAspGlnGlyAsnPheProLeuIleIleLysAsnLeuLysIleGluAspSerAspThrTyr
401 ATCTGTGAAGTGAAGACCAAGAGAGAGGAGTGCATTTGCTAGTGTCCGATTTGACTGCCAAGCTTGCACACCCAGCTGTTTCAGGGGAGAGGCTGACCC
TAGACACTTCACTCCTGCTTCCTCCCTCCAGCTTAACGATCACAAGCCCTAACCTGACCGGTGAGACTGTGGGTGAGACGAAGTCCCGCTCCGACTGGG
83 IleCysGluValGlnAspGlnGlyGlnGluValGlnLeuValPheGlyLeuThrAlaAsnSerAspThrHisLeuLeuGlnGlnGlyGlnSerLeuThrLeu
501 TGACCTTGAAGAGCCCCCTGCTGATAGCCCTTCACTGCAATGTAGAGAGTCCCAAGGGGTAAAAAACATACAGGGGGGAGAGACCTCTCCGTGTCTACCT
ACTGAACCTCTCGGGGAGCTATCGGGAGTCAAGCTTACCTCAGGTTCCCATTTTGTATGTCACCCCTCTCGGAGAGGACACAGCTCAG
117 ThrLeuGlnSerProProGlySerSerProSerValGlnCysArgSerProArgGlyLysAsnIleGlnGlyLysThrLeuSerValSerGlnLeu

FIG. 1B-1

bstXI
 aluI
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 bspI286
 banII
 bspI286
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 bstNI
 scrFI
 bstNI
 nlaIII
 mboII
 nheI
 aluI
 stuI
 haeIII
 mnlI
 150 GluLeuGlnaspSerGlyThrTrpThrCysThrValLeuGlnasnGlnLysValGluPheLysIleaspIleValValLeuAlaPheGlnLysAla
 601 GGAGCTCCAGATAGTGGCACCTGGACATGCACTGCTTCCAGAACCAAGAGTGAGTTCACAAATAGACATCGTGTGCTAGCTTCCAGAGAGCC
 CCTCGAGGTCTTATCACCGGTGACCTGTACGTGACAGACAGTCTGTCTTCCACCTCAAGTTTATCTGTAGCACACGATCGAAAGGCTCTCCG
 701 TCCAGCATAGTCTATAGAAAGGGGGAACAGGTGAGTCTCTCCCTCCACTCCGCTTTACAGTTGAAAAGCTGACGGGGCAGTGGCGAGCTGTGGTGGC
 AGTCGTATCAGATATTTCTTCCCTCCCTGTCCACCTCAAGAGAGAGGGGTGACCGAATGTCAACTTTGACTGCCCGTCACCGCTCGACACACCG
 183 SerSerIleValTyrLysLysGlnGluGlnValGluPheSerPheProLeuAlaPheThrValGluLysLeuThrGlySerGlyLnuLeuTrpGln
 mnlI
 aluI
 mnlI
 mnlI
 pflMI
 aluI
 hphI
 sau3AI
 dpuI
 mboII
 bstEII
 ecoO
 ddeI
 aluI
 scrFI
 bstNI
 sau96I
 nlaIV
 avaiI
 ppumI
 801 AGCGGAGAGGCTTCCCTCCCTCAAGTCTTGATCACCTTTGACCTGAAGAACAGGAAGTGTCTGTAAACGGGTTACCCAGAGCCTTAAGCTCCAGAT
 TCCGCTCTCCCGAAGAGAGAGGAGGAGTCAAGACCTAGTGAACCTGACTTCTGTCTTCCCTTCAAGACATTTGCCCCATGGGCTCTGGGATTCGAGGTCTA
 217 AlaglnArgAlaSerSerSerLysSerTrpIleThrPheaspLeuLysasnLysGlnValSerValLysArgValThrGlnaspProLysLeuGlnMet
 mnlI
 mnlI
 mnlI
 pflMI
 aluI
 hphI
 sau3AI
 dpuI
 mboII
 bstEII
 ecoO
 ddeI
 aluI
 scrFI
 bstNI
 sau96I
 nlaIV
 avaiI
 ppumI

sau96I
 avaiI
 ppuMI
 ecoO
 1201 CTGCGACAGGTCCTGCTGGAATCCACATCAAGTTCTGCCCCACATGGTCCACCCGAGCTTAATGCGGTAGTTATCAAGTTAATTGCTAACGCA
 GAGCCCTGCCAGACGACCTTAGTTGTTAGTTCACAGACGGGTGTAACAGTTGGGGCTCGAATTACGCCATCAATAGTGTCAATTACGATTGCCGT
 350 SerGlyGlnValLeuLeuGlnSerAsnIleYsValLeuProThrTyrPseThrProSerPheAsnAlaValValTyrHisSerOC*
 avaiI alvNI hinfi
 sau96I
 avaiI aluI
 mseI
 1301 GTCAGGCACCGGTATGAATCTAACAATGCGCTCATCGTCATCCCTCGGCACACCGTCACCCCTGATGCTGTAGGCATAGCCTTGTTATGCGGTAAGTGC
 CAGTCCGTGGCACATATCTTAGATTGTACCGAGTAGCAGTAGGAGCCGTGGCAGTGGAACCTACGACATCCGTATCCGAACCAATACGGCCATGACGG
 mnlI
 haeIII
 sau96I
 1401 GGGCTCTTGGCGGAT
 CCCGAGAACGCCCTA
 nlaIV
 bani
 hinPI
 hhaI
 mnlI
 nlaIV
 scriI
 bstNI
 foki
 bani
 hphi
 foki
 mspi
 hpaII
 hpaII
 mspi
 nciI
 mspi
 scriI
 sfanI
 rsaI
 mspi
 hpaII

FIG. 1C

FIG. 2A

| thai | scrFI |
|---------|---|
| alul | hstNI |
| hindIII | hpaII |
| hpaII | fnu4HI |
| sacl | nlaiV |
| dpnI | |
| afII | |
| mseI | |
| styI | MetGlyGlyThrAlaAlaArgLeuGly |
| 1 | TTGCAAGTCGCGCTTGGTGGTGTGATGGGCTAGTAGTCATATGGAATTCCAGAAACACACCCAGCAAGGCCATATACCCGCCCTGACGGCGGTCCAAACC |
| 1 | AAGCTTCAGCGCGCAACGACCAACTATCCCCGATCATCTATATCCCTTAAGTCTCTTTTGTGTGGTGCGCTTCCGGTATGGGGGGAAGTGCAGCCAGGTTGCG |
| 1 | alul hindIII hpaII sacl dpnI afII mseI hpaII fnu4HI nlaiV |

[illegible][illegible]

| | mbolI | mbolI | hinfI | nlaiV | fokI |
|-----|-----------------------------------|---------------------------------|--|-------------------------------------|------|
| 301 | TCCGAGAGAGAGCATATCCACTCGAATAA | CTCCAA | CCAGATTAAGATTCTCGGAATCAGAGGCTCTTCTTAAC | TAAGAGTTCATCCACAGCTGA | |
| | AGGGCTTCTTCTCGTATGTTAAGGTGACCTTTT | TGAGGTGGCTATTTCTTAAGACCTTTAGTCC | CAAGAAAGATTGATTTCCAGGTAGGTTGCACT | | |
| 76 | SerGlnIyIySerIleGlnpHehIstrpIySa | nSerIa | nGlnIleYsIleuclnIySa | nGlnIySerpHeIeuThrIyGlyProSerIySeIa | nS |

[illegible]

| | avali | mnlI | econI | alwNI | styI |
|--|---|------|-------|-------|------|
| 501 | AGTGAGGACCAGAGGAGGAGGTGCATTTGCTAGTGTTCGGAATTGACTGCGCAACTGTGACACCCGACTGCTTACAGGGGAGAGCCTGACCCCTGACCTTG | mnlI | bspMI | | |
| TCACCTCTCGGTCTTCCCTCCCTCCACGTTAACTCATCACAGCCTCAACTGACGGGTGACGAGAGTCCCCCTCTCGGACTCGGACTGGAGCTGGAAAC | | | | | |
| 143 | ValGluAspClnIySGluGluValAlnIeuIeuValPheGlyIeuThrAlaAsnSerAspThrHisIeuIeuGlnGlyGlnSerIeuThrIeuThrIeu | | | | |

SCIFI
POSTXI

701 AGGATAGTGGCACTTGACATGCACTGTCTTGCAGAACCAAGAGCTTGCACATTCGCTGCTTCCAGAGGCTCCAGAT
 TCCATACCTGGGAGCTGACGTGACAGAACGCTGCTTCCACCTCAAGTTTATCTGACAGCAGCATGCAAGGCTCTCCGAGGTGTA
 210 AspSerGlyThrTrpThrCysThrValLeuGlnAsnGlnLysValAlaGlnPheLysIleAspIleValValLeuAlaPheGlnLysAlaSerSerIle
 801 AGCTATATAAGAAAGAGGGGGAACAGCTGGAGTTCTCTTCCCACTCGCCCTTTACAGTTGAAAAAGCTGACCGGGCAGTGGCGAGCTGTGTCGACAGCGGAGC
 TCAGATATTCTTTCTCCCTGCTGTCACCTCAAGAGGAGGCTGAGCGGAATGTCACACTTTTCGATGCACTGCGCCGTACACCGCTCCGCTC
 243 ValTyrLysLysGlnGlyGlnGlnValGlnPheSerPheProLeuAlaPheThrValGlnLysLeuThrGlySerGlyGlnLeuTrpTrpGlnAlaGln
 901 AGGGCTTCTCTCCCAAGTCTTGACATGACCTTTGACCTGAGAGAACAGAGAGTGTCTGTAAACGGGTACCCAGACCTTAAGCTCCAGATGGGCAAGA
 TCCCGAAGGAGGAGGTTCAAGAACCTAGTGAACCTGACATCTTGTCTTCAACAGACATTTTGCCCAATGGGTCTGTGGATTCAGGCTACCCGTTCT
 276 ArgAlaSerSerSerLysSerTrpIleThrPheAspLeuLysAsnLysGlnValSerValLysArgValAlaThrGlnAspProLysLeuGlnMetGlyLysLys

1001 AGTCCCGCTCCACCTCACCCTGCCCCAGGCCCTTGCTCAGTATGCTGGCTCTGGAAACCTCACCCTGCCCCCTTGAAGCGAAACAGGAAAGTTGCATCA
 alui hphi scrii haeIII stuI sau96I
 mli econI bstNI ddeI hphi scrii haeIII
 310 TCGAGGCGGAGTGGAGTGGACGGGCTCCGGAACGAGTCATACGACCGAGACTTGGAGTGGACCGGGAACCTTCGCTTTGTCTTCAACGTA
 LeuProLeuH1sLeuThrLeuProGlnAlaLeuProGlnTyrAlaGlySerGlyAsnLeuThrLeuAlaLeuGlnAlaIySThrGlyLysLeuH1sGln
 1101 GGAAGTGAACCTGGTGGTGAAGAGCCACTCAGCTCCAGAAATTTGACCTGTGAGGTGTGGGAGCCACCTCCCTTAAGCTGATGCTGAGTTTGAAA
 bstNI hphi ddeI alui mli econI
 343 CCTCAGTTGGACCACTACTCTCGGTGAGTCGAGGTCTTTTAAACTGACACTCCACACCCCTGGGTGAGGGGATTCGACTACGACTCAAACTTT
 GluValAsnLeuValAlaMetArgAlaThrGlnLeuGlnLysAsnLeuThrCysGluValTrpGlyProThrSerProLysLeuMetLeuSerLeuLys

FIG. 2B-2

CD4



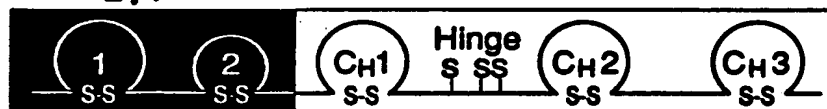
Immunoglobulin γ_1



Soluble rCD4



CD4₂ γ_1



CD4₄ γ_1



FIG. 3

FIG. 4A

1 GAATTCTGTCACTGCGCGGACACGCGCCGTATATTACTGTGCGAGAGACCACCTTTTGGCTATGTTACAGGAGCGTCCCGCTTGTGGATCGACCCCTGG
 2 CTTAAGACAGTGAAGCGCGCTGTGCGCGATATATGACACAGCTGTGGAAGAGGATACCATGTGCTCGGACAGGGGAGACCACTAGCTGGGACC
 3 ValThrAlaAlaAspThrAlaValTyrTyrCysAlaArgAlaThrPheCysLeuTyrPyrArgGluArgProProCysTyrPileAspProTyr
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FIG. 4B-1

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sau96I
nlaIV
mspi
sau3AI avariI mnlI
nlaIII hpaII ddel
bsphI scrFI mstII
mboII
mnlI styI mnlI dphI nciI eco8II nlaIII
501 TCCCTCTCCCCCAAAACCCAGAGACACCCCTCATGATCTCCCGACCCCTGAGTCACATGCCGTGGTGGAGCTGAGCCACGACCACTCCACTCGGTGCTTGGAGCTCCAGTT
AGGAGAGAGGGGGGTTTGGGTTCTCTGTGGAGTACTAGAGGGCCCTGGGGACTCCAGTACGACCACTCCACTCGGTGCTTGGAGCTCCAGTT
237 LeuPheProProlYsProlYsAspThrLeuMetIleSerArgThrProGluValThrcysValValValAspValSerHisGluAspProGluValLys

mboII
mnlI
rsai mnlI
fnu4HI mnlI rsai rsai rsai hgaI hphI
sacII
that
scrFI
ncII
mspi
hpaII
mnlI
601 GTTCAACTGGTACGTGAGACGGCGTGGAGGTGCATTAATGCCAAGACAAGCCCGGGAGGAGCAGTACAAACAGCAGTACCGGGGTGGTCAAGCTTCCAC
CAAGTTGACCATGCACCTGCCGCACTCCACGTAATTACGGTTCTGTTTCGGCCCTCTCTGCATGTGTGTCGATGGCCACACAGTCCAGAGGTG
270 PheAsnTrpTyrValAspGlyValGluValHisAsnAlaLysThrLysProArgGluGluGlnTyrAsnSerThrTyrArgValValSerValLeuThr

scrFI
econI bstNI
rsai
mnlI
tagI
701 GTCCGTGCACACGAGACTGGCTGAATGCCAAGAGTACCAAGTCTCCACAAGCCCTCCAGCCCCCATGAGAGAAACCATCTCCAAAGCCAAAG
CAGGACGTGGCTGTGACCGCACTTACCGTTCCTCATGTTCACGTTCCAGAGGCTGTCGGAGAGGTGCGGGGTAGCTCTTTGGTAGAGGTTTCGCTTC
303 ValLeuHisGlnAspTrpLeuAsnGlyLysGlnTyrLysCysLysValSerAsnLysAlaLeuProAlaProIleGlnLysThrIleSerLysAlaLysGly

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scrFI
ncII
mspi
hpall
smal
scrFI
ncII
aval
fnu4HI
rsal
fokI
fokI
alul
bscFI
bstNI
bspMI
bscFI
bstNI
801 GGCAGCCCCGAGAACCACAGGTGTACACCCCTGCCCCCATCCCCGGGATGAGCTGACCAAGAACCAGCTCAGCCCTGACCTGCTCAAAAGCCTTATCC
bvi avai
337 GlnProArgGluProGlnValTyrThrLeuProProSerArgAspGluLeuThrLysAsnGlnValSerLeuThrCysLeuValIysGlyPheTyrPro
fnu4HI
mspi
hpall
901 CAGGACATCGCCGCTGAGTGGAGAGACAATGGGACAGCCGAGAGAGAACAACTACAGACCAGCCCTCCCGCTGCTGACTCCGACGGCTCCCTTCTCTCTAC
GTCGCTGAGCGGCACCTCACCCTCTCGTTACCCCGCGCTCTGTTGATGTTCTGCTGGAGGCGACGACCTGAGGCTGCCGAGAGAGAGATG
370 SerAspIleAlaValGluTrpGlnSerAsnGlyGlnProGluAsnAsnTyrIleThrThrProValLeuAspSerAspGlySerPheLeuTyr
mspi
bvi
fnu4HI
hpaII
mnlI
pleI
hinfI
nlaIV
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mnlI
1001 AGCAAGCTCACCGTGGACACAAGAGGAGGTGGCAGCAGGGGGAACGCTTCTCAAGCTCCGCTGATGATGAGGCTCTGCACAACCACTACACGAGAGAGCC
alul
bspMI
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mboII
mnlI
403 SerLysLeuThrValAspLysSerArgTrpGlnGlnLysAsnValPheSerCysSerValMetHisGlnAlaLeuHisAsnHisTyrThrGlnLysSerLeu
hphI
alul
bvi
fnu4HI
xmiI
mboII
nlaIII
nsII
avalIII
scrFI
ncII
mspi
hpall
haeIII
xmaIII
eaeI
1101 TCTCCCTGTCTCCGGGTAAATGAGTCCGACGCGCC
AGAGGACAGAGGCCCATTTACTCACGCTGCCGCC
437 SerLeuSerProGlyLysOP*

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FIG. 4B-2

[illegible]